Figure 1.

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ln-Se	_	0.51 0.26 0.16					
al-G	Val	GUU 0.51 GUA 0.26 GUG 0.16 GUC 0.07	125				
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Lys-1	Leu	CUG 0.83 CUC 0.07 CUU 0.04 UUG 0.03 UUA 0.02	ın				
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(a) Nucleotide sequences for Ala-Ile-Lys-Leu-Val-Gln-Ser-Pro and showing codon frequencies in relation to usage in enteric bacterial genes	Ala	GCU 0.35 GCA 0.28 GCG 0.26 GCC 0.10	178	8 Pro	CCG 0.77 CCA 0.15 CCU 0.08 CCC 0.00		
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1-Phe-1	ų	0.37 0.20 0.03 0.03	ત	0.35 0.28 0.10		
ly-Asn	Ser	UCC UCC AGC UCG AGU UCA	Ala	600 600 600 600 50		
Asn-G	Gln	0.86 0.14	Ala	0.35 0.28 0.26 0.10		
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r Ala- usage	Lys	AAA 0.74 AAG 0.26	Gly	0.59		
es for	j.	AAA AAG	ָם מ	660 660 664 664		
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(b) Nucleotide sequences for Ala-Ile-Lys-Leu-Val-Gln-Ser-Pro-Asn-Gly-Asn-Phe-Ala-Ala-Ser frequencies in relation to usage in enteric bacterial genes	Ala		6 8 E	000 000 000 401	15 Ser	000 000 000 000 000 119

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-Thr- genes.								
-Lys-Leu-Val-Gln-Ser-Pro-Asn-Gly-Asn-Phe-Ala-Ala-Ser-Phe-Val-Leu-Asp-Gly-Thr- and showing codon frequencies in relation to usage in enteric bacterial gene Leu Val Gln Ser								
eu-As bact								
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n-Phe-; elation Ser		0.02	Ala		0.10	Thr	0.55	
y-Asn in re	UCC UCC AGC UCG	158	Ø	900 800 800	ರೆ 50	Ħ	ACC ACC ACG ACG	338
sn-Gl cies n	CAG 0.86		ď	0.35	0.10	≻ ,	0.59 0.38 0.02 0.00	
Pro-Ass equenc: Gln	CAG	230	Ala		34 34	Gly	660 660 666 668	240
-Ser- lon fr	0.51 0.26 0.16 0.07		0	0.76		0.	0.67	
il-Gln ig cod Val	GUA GUA GUG GUG GUC	125	Phe	200	34	Asp	GAC (GAU (161
Lys-Leu-Val- and showing Leu	0.83 0.07 0.03 0.02	0.00		0.94			0.03	000
Lys-I and s Leu		CUA 0.	Asn	AAC 0	88	Leu		
a-Ile r-Tyr								
or Ala ys-Ty: Lys	AAA 0.74 AAG 0.26	269	Gly	GGU 0.59 GGC 0.38 GGG 0.02	GGA 0.00 148	Val	GUU 0.51 GUA 0.26 GUG 0.16 GUC 0.07	167
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equen -Lys-: Ile	0.83 0.00 0.00	0	Asn	AAC 0.94 AAU 0.06	6	Phe	C 0.76 U 0.24	vo
ide se e-Phe	AUC AUU AUA	260	14		396	21	000	216
ucleot: Trp-Ile	0.35 0.28 0.26 0.10		B -	0.77		Ser -		0.03
(c)Nucleotide sequences for Ala-Ile Lys-Trp-Ile-Phe-Lys-Ser-Lys-Tyr-Tyr Ala Ile Lys	900 900 900 900 900	178	ထမ်	8 9 9	401	15 S. S.	UCC UCC AGC	HGG 113
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Lys	AAA 0.74 AAG 0.26	o			
Ser	UCC 0.37 UCU 0.34 AGC 0.20 UCG 0.04	154 154			
Lys	AAA 0.74 AAG 0.26	152			
Phe	UUC 0.76 UUU 0.24	154			
IJe	AUC 0.83 AUU 0.17 AUA 0.00	173			
£	UGG 1.00	467	Tyr	UAC 0.75	0
22 - 28	Lys AAA 0.74 AAG 0.26	467	29 - 30 Tyr	UAC 0.75 UAU 0.25	0